

Please replace the three paragraphs beginning at page 5, line 21, and ending at page 6, line 5, with the following rewritten paragraphs:

--In one embodiment, a peptide according to (i) (a) above has a sequence selected from:

(a1) Leu Lys Thr Pro Arg Val (SEQ ID NO:1)

(a2) Lys Thr Pro Arg Val Thr (SEQ ID NO:2)

(A) Asn Thr Leu Lys Thr Pro Arg Val Gly Gly (SEQ ID NO:3)

In another embodiment, a peptide according to (i) (b) above has a sequence selected from:

(b1) Lys Asp Lys Ala Thr Phe (residues 1-6 of SEQ ID NO:4)

(B) Lys Asp Lys Ala Thr Phe Gly Thr His Asp Gly (SEQ ID NO:4)

In a further embodiment, a peptide according to (i) (c) above has a sequence selected from:

(c1) Thr Leu Arg Val Tyr Lys (residues 3-8 of SEQ ID NO:7)

(c2) Thr Lys Leu Arg Val Tyr (SEQ ID NO:5)

(c3) Thr Leu Leu Arg Val Tyr (SEQ ID NO:6)

(C) Cys Ala Thr Leu Arg Val Tyr Lys Gly Gly (SEQ ID NO:7) --

Please replace the paragraph beginning at page 13, line 4, with the following rewritten paragraph:

--Examples of divalent peptides with peptides derived from mono-peptides A-C of the invention and irrelevant peptide D are as follows:

A: Asn Thr Leu Lys Thr Pro Arg Val Gly Gly X Ala (SEQ ID NO:8)

B: Lys Asp Lys Ala Thr Phe Gly Thr His Asp Gly Gly X Ala (SEQ

ID NO:9)

C: Cys Ala Thr Leu Arg Val Tyr Lys Gly Gly Gly X Ala (SEQ ID NO:10)

D: Pro Val Arg Ser Pro His Gln Ser Tyr Pro Gly Gly Gly X Ala (SEQ ID NO:11)

wherein X=FmocLys(Fmoc)-OH--

Please replace Table 1 on page 22 with the following rewritten Table 1:

--Table 1

Binding of anti- β 2GPI mAbs to phage isolated from an hexapeptide epitope library

Antibody	Epitope sequence*	SEQ ID NO:	Binding O.D.	Phages identified, no.
ILA-1	L K T P R V	8 (res. 3-8)	975 \pm 72	24
	K T P R V T	12	1201 \pm 142	18
ILA-3	K D K A T F	13	1178 \pm 101	15
G-3	T L R V Y K	14	791 \pm 33	12
	T K L R V Y	15	954 \pm 62	7
	T L L R V Y	16	869 \pm 71	16

Please replace the two paragraphs beginning at page 23, line 1, and ending at page 23, line 11, with the following rewritten paragraphs:

--ILA-1 mAb detected the sequence KTPRV (residues 4-8 of SEQ ID NO:8) that appeared in 42 clones, wherefrom 24 clones showed the sequence LKTPRV (residues 3-8 of SEQ ID NO:8) which represents a mimotope located between domain I/II of the β 2GPI molecule as LK(C)TPRV (SEQ ID NO:17) on the native form of the β 2GPI . The other 18 clones showed the